Supplementary Material

Alterations in Morphology and Adult Neurogenesisin the Dentate Gyrus of *Patched1* Heterozygous Mice

Antonelli F., Casciati A., Tanori M., Tanno B., Linares-Vidal M.V., Serra N., Bellés M., Pannicelli A., Saran A., Pazzaglia S.*

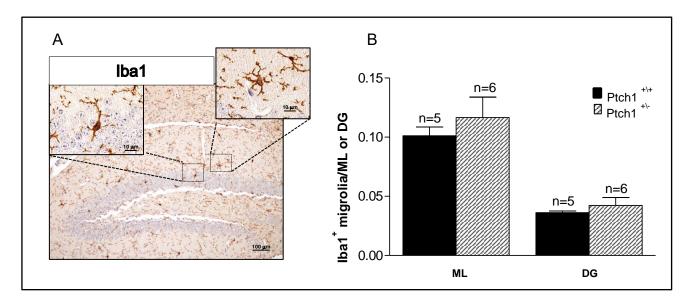


FIGURE S-1: Shh pathway deregulation does not affect microglia. (A) Immunostaining and (B) quantification of Iba1 positive microglia cells in the ML and DG of the hippocampus of 8-month old mice. Images, 10x magnification, scale bar = 100 μm ; 100x magnification, scale bar = 10 μm . The number of mice used per test is indicated in the graph (n). Statistical analysis was determined using a two-tailed Student's t-test for comparison between pairs of means.

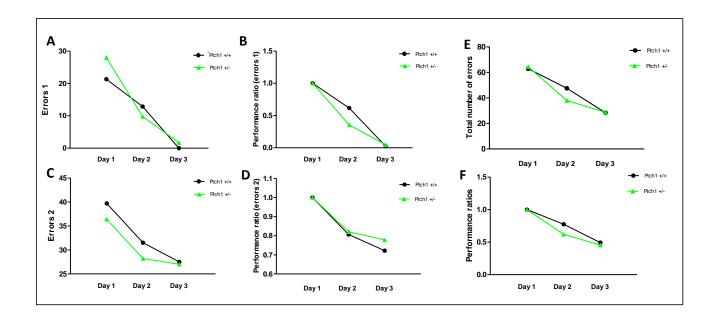


FIGURE S-2: Lack of effect of Shh signaling on spatial working memory using Radial Arm Maze (RAM) test at 4 months of age. $Ptch1^{+/+}$ (n=10) and $Ptch1^{+/-}$ (n=10) mice submitted to RAM test. (A) Errors type 1 (animal visits an arm and did not eat the reward). (B) Performance ratio 1 (errors type 1 standardized to day 1). (C) Errors type 2 (animal visits the same arm more than once during a single test session). (D) Performance ratio 2 (errors type 2 standardized to day 1). (E) Total number or errors (Errors 1 + Errors 2). (F) Performance ratios (total number of errors standardized to day 1). Data are given as the mean \pm standard error (SEM). Statistical significance for all tests was established at p<0.05.

TABLE S-1: Primer Sequences used for Real-Time Quantitative PCR.

| Gene | Forward primer | Reverse primer |
|-----------|----------------------------------|---------------------------------|
| Gli1 | 5' – GAGGACCTGGAGAGAGAGAGAA - 3' | 5' - CCAGCGGCAGTCTGTCTCA - 3' |
| TLX | 5' - CGATTAGACGCCACTGAA - 3' | 5' – GGTATCTGGTATGAATGTAGC – 3' |
| Cyclin D1 | 5' - GCAAGCATGCACAGACCTT- 3' | 5' – GTTGTGCGGTAGCAGGAGA – 3' |
| | | |

TABLE S-2: Expression changes in neurogenesis-related genes in $Ptch1^{+/-}$ compared to WT mice at 8 months of age. Significantly up- and down-regulated genes are indicated in red and blue, respectively. A cut-off of 1.3-fold in change of gene expression was applied.

| Gene Symbol | Gene Title | Fold Regulation | p-value |
|-------------|---|-------------------|----------|
| Ache | Acetylcholinesterase | 1,0193 | 0,801363 |
| Adora1 | Adenosine A1 receptor | -1,011 | 0,789168 |
| Adora2a | Adenosine A2a receptor | 1,7751 | 0,363635 |
| Alk | Anaplastic lymphoma kinase | 1,0263 | 0,84238 |
| Apbb1 | Amyloid beta (A4) precursor protein-binding, family B, member 1 | 1,4778 | 0,034366 |
| Apoe | Apolipoprotein E | -1,041 | 0,699042 |
| Арр | Amyloid beta (A4) precursor protein | 2,1096 | 0,065162 |
| Artn | Artemin | 1,0093 | 0,949386 |
| Ascl1 | Achaete-scute complex homolog 1 (Drosophila) | -1,0559 | 0,659811 |
| Bcl2 | B-cell leukemia/lymphoma2 | -1,3134 | 0,114788 |
| Bdnf | Brain derived neurotrophic factor | 1,0287 | 0,649982 |
| Bmp2 | Bone morphogenetic protein 2 | 1,3263 | 0,966947 |
| Bmp4 | Bone morphogenetic protein 4 | 1,0838 | 0,396875 |
| Bmp8b | Bone morphogenetic protein 8b | 1,8556 | 0,054777 |
| Cdk5r1 | Cyclin-dependent kinase 5, regulatory subunit 1 (p35) | 1,4105 | 0,000718 |
| Cdk5rap2 | CDK5 regulatory subunit associated protein 2 | 1,4586 | 0,007103 |
| Chrm2 | Cholinergic receptor, muscarinic 2, cardiac | -1,5223 | 0,026664 |
| Creb1 | CAMP responsive element binding protein 1 | -1,2056 | 0,089482 |
| Cxcl1 | Chemokine (C-X-C motif) ligand 1 | -21,214 | 0,016508 |
| Dcx | Doublecortin | 1,3752 | 0,066337 |
| Dlg4 | Discs, large homolog 4 (Drosophila) | 1,3581 | 0,043013 |
| DII1 | Delta-like 1 (Drosophila) | 1,2717 | 0,351702 |
| Drd2 | Dopamine receptor D2 | 1,3107 | 0,590797 |
| Dvl3 | Dishevelled 3, dsh homolog (Drosophila) | 1,0116 | 0,930631 |
| Efnb1 | Ephrin B1 | 1,2053 | 0,196411 |
| Egf | Epidermal growth factor | -1,0915 | 0,386507 |
| Ep300 | E1A binding protein p300 | 1,4764 | 0,123412 |
| Erbb2 | V-erb-b2 erythroblastic leukemia viral oncogene homolog 2, neuro/glioblastoma derived oncogene homolog (avian) | 2,7432 | 0,032266 |
| Fgf2 | Fibroblast growth factor 2 | 1,0482 | 0,639898 |
| Flna | Filamin, alpha | 1,8684 | 0,00437 |
| Gdnf | Glial cell line derived neurotrophic factor | 1,792 | 0,199044 |
| Gpi1 | Glucose phosphate isomerase 1 | 1,1516 | 0,044462 |
| Grin1 | Glutamate receptor, ionotropic, NMDA1 (zeta 1) | 1,652 | 0,003729 |
| Hdac4 | Histone deacetylase 4 | -1,413 | 0,091123 |
| Hes1 | Hairy and enhancer of split 1 (Drosophila) | -1,1447 | 0,072381 |
| Hey1 | Hairy/enhancer-of-split related with YRPW motif 1 | 1,3956 | 0,003629 |
| Hey2 | Hairy/enhancer-of-split related with YRPW motif 2 | -1,4647 | 0,010087 |
| Heyl | Hairy/enhancer-of-split related with YRPW motif-like | 1,2233 | 0,372172 |
| II3 | Interleukin 3 | -1,2151 | 0,518317 |
| Mdk | Midkine | 1,0989 | 0,281474 |
| Mef2c | Myocyte enhancer factor 2C | 2,2004 | 0,023552 |
| Kmt2a | Myeloid/lymphoid or mixed-lineage leukemia 1 | 2,6058 | 0,023332 |
| Map2 | Microtubule-associated protein 2 | 1,0846 | 0,377087 |
| Ndn | Necdin | -1,0582 | 0,649536 |
| Ndp | Norrie disease (pseudoglioma) (human) | -1,0362 | 0,317091 |
| Neurod1 | Neurogenic differentiation 1 | -1,2303 | 0,062632 |
| | | | |
| Neurog1 | Neurogenin 1 | -3,2924 7,4928 | 0,044035 |
| Neurog2 | Neurogenin 2 Neurofibromatosis 1 | 7,4938 1,4285 | 0,0217 |

| Gene Symbol | Gene Title | Fold Regulation | p-value |
|-------------|---|-----------------|----------|
| Nog | Noggin | 1,1322 | 0,340875 |
| Notch1 | Notch gene homolog 1 (Drosophila) | 1,7943 | 0,076791 |
| Notch2 | Notch gene homolog 2 (Drosophila) | 2,1685 | 0,012814 |
| Nr2e3 | Nuclear receptor subfamily 2, group E, member 3 | 1,4149 | 0,134383 |
| Nrcam | Neuron-glia-CAM-related cell adhesion molecule | -1,3396 | 0,051273 |
| Nrg1 | Neuregulin 1 | 1,0164 | 0,796512 |
| Nrp1 | Neuropilin 1 | -1,6076 | 0,006221 |
| Nrp2 | Neuropilin 2 | -1,0567 | 0,587594 |
| Ntf3 | Neurotrophin 3 | 1,6855 | 0,004428 |
| Ntn1 | Netrin 1 | 1,5268 | 0,099107 |
| Tenm1 | Odd Oz/ten-m homolog 1 (Drosophila) | -1,1639 | 0,476195 |
| Olig2 | Oligodendrocyte transcription factor 2 | -1,5311 | 0,08057 |
| | Platelet-activating factor acetylhydrolase, isoform 1b, | | |
| Pafah1b1 | subunit 1 | 1,0365 | 0,644714 |
| Pard3 | Par-3 (partitioning defective 3) homolog (C. elegans) | 1,372 | 0,006458 |
| Pax3 | Paired box gene 3 | -1,2866 | 0,490475 |
| Pax5 | Paired box gene 5 | 2,2816 | 0,08455 |
| Pax6 | Paired box gene 6 | 1,3846 | 0,039007 |
| Pou3f3 | POU domain, class 3, transcription factor 3 | 1,4539 | 0,156044 |
| Pou4f1 | POU domain, class 4, transcription factor 1 | 1,2213 | 0,416185 |
| Ptn | Pleiotrophin | -1,5426 | 0,002388 |
| Rac1 | RAS-related C3 botulinum substrate 1 | -1,147 | 0,022048 |
| Robo1 | Roundabout homolog 1 (Drosophila) | -1,2136 | 0,030484 |
| Rtn4 | Reticulon 4 | -1,2882 | 0,043584 |
| S100a6 | S100 calcium binding protein A6 (calcyclin) | -1,0518 | 0,537386 |
| S100b | S100 protein, beta polypeptide, neural | 1,0582 | 0,075549 |
| Shh | Sonic hedgehog | -1,6562 | 0,038125 |
| Slit2 | Slit homolog 2 (Drosophila) | 1,062 | 0,809289 |
| Sod1 | Superoxide dismutase 1, soluble | -1,206 | 0,002587 |
| Sox2 | SRY-box containing gene 2 | -1,0306 | 0,248389 |
| Sox3 | SRY-box containing gene 3 | 1,629 | 0,15878 |
| Stat3 | Signal transducer and activator of transcription 3 | 1,1534 | 0,126858 |
| Tgfb1 | Transforming growth factor, beta 1 | -1,0847 | 0,454668 |
| Th | Tyrosine hydroxylase | ND | ND |
| Tnr | Tenascin R | 1,1983 | 0,039991 |
| Vegfa | Vascular endothelial growth factor A | -1,0553 | 0,777697 |